

GenCore version 5.1.4 p5 4578
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OM protein - protein search, using sw model

Run on: March 17, 2003, 16:34:47 ; Search time 39 Seconds
(without alignments)
888.337 Million cell updates/sec

Title: US-09-840-243B-11

Perfect score: 1341
Sequence: 1 MELTQPAEDLITQOTPAE.....VIENHILKLFQSNLVPADPE 260

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-Processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

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1: A.GeneSeq_101002.1*
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3: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA1981.DAT:*
4: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA1982.DAT:*
5: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA1983.DAT:*
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14: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA1992.DAT:*
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16: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA1994.DAT:*
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22: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA2000.DAT:*
23: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA2001.DAT:*
24: /SIDSI/gcgdata/geneSeq/geneSeq-emb1/AA2002.DAT:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1341	100.0	260	21	AAV79411 Human MHC class II
2	1341	100.0	260	21	AAV58539 Human ankyrin repeat fami
3	1113.5	83.0	269	21	AAV59590 Ankyrin repeat anti
4	625	46.6	229	23	ABP41881 Human ovarian anti
5	621.5	46.3	313	22	AA66309 Human ankyrin-like
6	621.5	46.3	313	22	AA66309 Human ankyrin-like
7	607.5	45.3	263	22	AAU20646 Human secreted pro
8	495	36.9	152	22	AB550161 Human transcriptio
9	421.5	31.4	84	21	AA601584 Human secreted pro
10	393	29.3	119	22	AAU20665 Human secreted pro

11	364	27.1	105	22	AAU20558 Human secreted pro
12	299	22.3	81	23	ABB97342 Novel human protei
13	274	20.4	234	22	ABB61859 Drosophila melanog
14	257	19.2	49	22	ABB43550 Peptide #11056 enc
15	257	19.2	49	22	AA621225 Peptide #7659 enco
16	257	19.2	49	23	ABG45308 Human peptide enco
17	212	15.8	705	22	AA675604 Human colon cancer
18	210.5	15.7	1762	22	AAU96841 Rat Kidins220 prot
19	210.5	15.5	2443	22	ABB60521 Drosophila melanog
20	207.5	15.5	166	23	ABB78585 3 ankyrin repeat m
21	207.5	15.5	1715	22	AA638993 Human polypeptide
22	207.5	15.5	1715	22	AA638925 Human Kidins220 pr
23	207.5	15.5	1715	22	AAU96840 Rat Kidins220 prot
24	206.5	15.4	1763	23	AAU80244 Drosophila melanog
25	205.5	15.3	342	22	ABB59641 Ankyrin protein fr
26	205	15.3	348	19	AAW76075 D. immitis ankyrin
27	205	15.3	348	19	AAW76775 D. immitis ankyrin
28	205	15.3	348	21	AA615588 Ankyrin protein se
29	205	15.3	348	23	AAW76068 Full length ankyri
30	205	15.3	1745	19	AAW76068 D. immitis ankyrin
31	205	15.3	1745	19	AAW76068 D. immitis ankyrin
32	205	15.3	1745	21	AA615589 Ankyrin protein se
33	205	15.3	1745	23	AAO21368 Human ORF ORF2052
34	204.5	15.2	978	22	AA642288 Human protein kin
35	202.5	15.1	765	22	AAW79160 Human protein, SEQ
36	202.5	15.1	1872	22	AAW79160 Drosophila melanog
37	199	14.8	1498	22	ABB64857 Human cancer cell
38	198	14.7	627	23	AAO17136 Human death domain
39	196.5	14.7	551	22	AAE01035 Novel human diagno
40	196	14.6	1377	22	ABG08072 Arabidopsis thalia
41	195.5	14.6	456	21	AA612893 Arabidopsis thalia
42	195.5	14.6	456	21	AA627402 Human cell growth
43	195	14.5	435	22	AA666710 Human protein sequ
44	195	14.5	435	22	AA693879 Human protein sequ
45	193	14.4	4274	22	ABG00972 Novel human diagno

ALIGNMENTS

RESULT 1				
AAV79411	AAV79411 standard; Protein; 260 AA.			
AAV79411;	(first entry)			
01-AUG-2000	(first entry)			
Human MHC class II gene transcription factor RFXANK.				
RFXANK; HSRFXANK; human; transcription factor; MHC class II;				
chromosome 19p12; immunosuppressive; immunomodulator;				
antiinflammatory; antidiabetic; antiarthritic; therapy;				
inflammation; autoimmune diseases; transplant rejection;				
insulin dependent diabetes; multiple sclerosis;				
lupus erythematosus; rheumatoid arthritis; immunodeficiency.				
Homo sapiens.				
Location/Qualifiers				
Region	/note="ankyrin repeat region 1"			
Region	/note="ankyrin repeat region 2"			
Region	/note="ankyrin repeat region 3"			
EP995798-A1.				
26-APR-2000.				
24-OCT-1998;	98EP-0120085.			

PR 24-OCT-1998; 98EP-0120085.
 XX (NOVI-) NOVIMMUNE SA.
 XX Mastermak K, Reith W, Mach B;
 PI WPI, 2000-294958/26.
 DR N-PSDB; AA294868.
 XX Novel isolated transcription factor, RFXANK, useful for treating MHC
 PT class II deficiency and autoimmune disorders, e.g. insulin dependent
 PT diabetes and multiple sclerosis, restores the functional transcription
 PT of MHC class II genes
 XX
 XX Claim 1; Fig 3; 48pp; English.
 XX
 XX The present sequence is that of human RFXANK, a novel transcription
 CC factor that is a subunit of the RFX heterotrimeric transcription
 CC complex that binds to the conserved X box motif of all MHC class II
 CC gene promoters. The RFXANK gene is mutated in complementation group
 CC B MHC II deficiency patients. Mutations identified in patients
 CC include aberrant splicing and short deletions in exon 6. The
 CC invention provides inhibitors of RFXANK including antibodies, single
 CC chain antibodies, dominant negative mutants, antisense molecules
 CC and ribozymes. The inhibitors may be used in therapy or prevention
 CC of diseases associated with aberrant expression of MHC class II
 CC genes and/or as immunosuppressive agents, e.g. to treat
 CC inflammation, autoimmune diseases or rejection of transplanted
 CC organs, insulin dependent diabetes, multiple sclerosis, lupus
 CC erythematosus and rheumatoid arthritis. The compositions may also
 CC be used to treat the autosomal recessive disease MHC class II
 CC deficiency. Since RFXANK does not play any other major role in the
 CC transcriptional control of genes other than MHC class II genes, its
 CC inhibitors are devoid of other undesirable inhibitory effects.
 XX
 XX Sequence 260 AA:
 SQ
 Query Match 100.0%; Score 1341; DB 21; Length 260;
 Best Local Similarity 100.0%; Pred. No. 2e-128;
 Matches 260; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MELTOPAEDLIQTQOTPASELGDPDEGEAAGSDTVVLSLFCPTPEPVNPEPDASVSS 60
 DB 1 MELTOPAEDLIQTQOTPASELGDPDEGEAAGSDTVVLSLFCPTPEPVNPEPDASVSS 60
 QY 61 PQAAGSLKSTLTITNRQGNVSALPATLDSLSIHQAAGELDQKHLRKGNDLVNKP 120
 DB 61 PQAAGSLKSTLTITNRQGNVSALPATLDSLSIHQAAGELDQKHLRKGNDLVNKP 120
 QY 121 DERGFPLIMASAFGEIETVRFLEMGADPHILAKERSALSASTGTYDVGILLERD 180
 DB 121 DERGFPLIMASAFGEIETVRFLEMGADPHILAKERSALSASTGTYDVGILLERD 180
 QY 121 DERGFPLIMASAFGEIETVRFLEMGADPHILAKERSALSASTGTYDVGILLERD 180
 DB 121 DERGFPLIMASAFGEIETVRFLEMGADPHILAKERSALSASTGTYDVGILLERD 180
 QY 181 VDINIYDMNGSTPLLYAVRGNHKVCVEALLARGADLTTEADSGTYPMDLAVAGRYKVOQ 240
 DB 181 VDINIYDMNGSTPLLYAVRGNHKVCVEALLARGADLTTEADSGTYPMDLAVAGRYKVOQ 240
 QY 241 VIENHILKLFQSNLVPADE 260
 DB 241 VIENHILKLFQSNLVPADE 260
 RESULT 2
 AA59539
 ID AA59539 standard; Protein; 260 AA.
 XX AA59539;
 AC
 XX 03-APR-2000 (first entry)
 DT
 XX Human ankyrin family protein, ANFP.
 DE Human ankyrin family protein; ANFP; autoimmune disorder; inflammation;
 XX Human; ankyrin family protein; ANFP; autoimmune disorder; inflammation;

KW atherosclerosis; inflammatory disorder; proliferative disorder; AIDS;
 KW vesicle-traffic disorder; allergy; amyloidosis; anaemia; asthma;
 KW bronchitis; Crohn's disease; atopic dermatitis; diabetes mellitus;
 KW irritable bowel syndrome; osteoporosis; rheumatoid arthritis; cirrhosis;
 KW hepatitis; ulcerative colitis; cancer; hypercholesterolaemia; therapy;
 KW diagnosis.
 XX
 XX Homo sapiens.
 OS
 XX US598963-A.
 XX
 XX 23-NOV-1999.
 PD
 XX 14-OCT-1998; 98US-0172977.
 PF
 XX 14-OCT-1998; 98US-0172977.
 XX
 XX 14-OCT-1998; 98US-0172977.
 XX
 XX (INCY-) INCYTE PHARM INC.
 XX
 XX Tang YT, Corley NC, Yue H, Guegler KJ;
 PI WPI: 2000-095634/08.
 DR N-PSDB; AA249052.
 XX
 XX polynucleotide sequence encoding a human ankyrin family protein useful
 PT for diagnosis or treatment of autoimmune, inflammatory, proliferative
 PT and vesicle-traffic disorders
 XX
 XX Claim 1; Fig 1; 34pp; English.
 XX
 XX This sequence is the human ankyrin family protein, ANFP, of the
 CC invention. Host cells containing an expression vector containing the
 CC polynucleotide sequence can be cultured to produce ANFP, which can be
 CC used for diagnosis or treatment of autoimmune, inflammatory,
 CC proliferative and vesicle-traffic disorders. Disorders which can be
 CC treated include acquired immune deficiency syndrome (AIDS), allergies,
 CC amyloidosis, anaemia, asthma, atherosclerosis, bronchitis, Crohn's
 CC disease, atopic dermatitis, diabetes mellitus, irritable bowel syndrome,
 CC myocardial or pericardial inflammation, osteoporosis, rheumatoid
 CC arthritis, cirrhosis, hepatitis, ulcerative colitis, cancer and
 CC hypercholesterolaemia. The polynucleotide sequences can also be used as a
 CC hybridisation probe to detect ANFP-encoding polynucleotides in biological
 CC samples. Purified ANFP can be used to produce antibodies or to screen
 CC libraries of pharmaceutical agents to find agents that specifically bind
 CC ANFP. The DNA and its antisense sequence can be used in therapeutic
 CC compositions e.g. to regulate gene function. The DNA sequence can be used
 CC for diagnostic purposes to detect and quantitate gene expression in
 CC biopsied tissues and to indicate the absence, presence and excess
 CC expression of ANFP and monitor its levels during therapeutic
 CC intervention.
 CC
 XX Sequence 260 AA;
 SQ
 Query Match 100.0%; Score 1341; DB 21; Length 260;
 Best Local Similarity 100.0%; Pred. No. 2e-128;
 Matches 260; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MELTOPAEDLIQTQOTPASELGDPDEGEAAGSDTVVLSLFCPTPEPVNPEPDASVSS 60
 DB 1 MELTOPAEDLIQTQOTPASELGDPDEGEAAGSDTVVLSLFCPTPEPVNPEPDASVSS 60
 QY 61 PQAAGSLKSTLTITNRQGNVSALPATLDSLSIHQAAGELDQKHLRKGNDLVNKP 120
 DB 61 PQAAGSLKSTLTITNRQGNVSALPATLDSLSIHQAAGELDQKHLRKGNDLVNKP 120
 QY 121 DERGFPLIMASAFGEIETVRFLEMGADPHILAKERSALSASTGTYDVGILLERD 180
 DB 121 DERGFPLIMASAFGEIETVRFLEMGADPHILAKERSALSASTGTYDVGILLERD 180
 QY 181 VDINIYDMNGSTPLLYAVRGNHKVCVEALLARGADLTTEADSGTYPMDLAVAGRYKVOQ 240
 DB 181 VDINIYDMNGSTPLLYAVRGNHKVCVEALLARGADLTTEADSGTYPMDLAVAGRYKVOQ 240

OY 241 VIENNHLKLFQSNLVPADPE 260
 DB 241 VIENNHLKLFQSNLVPADPE 260

RESULT 3

ID AAY59590 standard; Protein; 269 AA.

AC AAY59590;

DT 03-APR-2000 (first entry)

DE Ankyrin repeat protein TVL-1.

KM Ankyrin repeat protein; TVL-1; TNF: tumour necrosis factor; apoptosis;
 cell cycle regulation; apoptotic cell death; cell proliferation.

OS Mus sp.

PN WO967269-A1.

PF 29-DEC-1999.

PR 24-JUN-1999; 99WO-US14353.

PR 24-JUN-1998; 98US-0090742.

(UYDE-) UNIV JEFFERSON THOMAS.

TS Ichl1s PN, Makris A;

WPI: 2000-106276/09.

DR N-PSDB; AAZ49069.

PT Novel isolated nucleic acid useful for biological screens to identify
 PT therapeutic agents involved in regulation of cell cycle progression and
 PT apoptotic cell death -

PS Claim 12; Fig 2C; 112pp; English.

CC This sequence is the ankyrin repeat protein, TVL-1 of the invention.
 CC The protein contains ankyrin repeat domains and is capable of promoting
 CC TNF (tumour necrosis factor) induced apoptosis. The nucleic acid
 CC molecules, proteins and antibodies are useful as targets for screening
 CC therapeutic agents that regulate cell cycle progression and apoptotic
 CC cell death, especially useful for identification, detection and/or
 CC regulation of complex signalling events that regulate cell cycle
 CC progression and apoptotic cell death. The TVL-1 molecules of the
 CC invention can also be used as a research tool and will facilitate the
 CC elucidation of the mechanistic action of the novel genetic and protein
 CC interactions involved in the control of cellular proliferation and
 CC apoptosis.

XX Sequence 269 AA;

Query Match 83.0%; Score 1113.5; DB 21; Length 269;

Best Local Similarity 81.9%; Pred. No. 3.6e-105; Indels 11; Gaps 2;

Matches 221; Conservative 15; Mismatches 23; Indels 11; Gaps 2;

OY 1 MELTPAEDLITQOTPAELGDEPDEBEADGSDTVVLSPCTPEPVPNEPPASVSS 60

DB 1 MEPTQVAENLVNQPVPVDEPDDEPDESDTVVLSFCTPPAVNPEADASAS 60

OY 61 PQAGSSLKHTTTNNRQGNVSALPATLDSISHQLAAGSLDQKHLRK----- 112

DB 61 LQ-GSFLKHSSTLTNNRQGNVSALPATLDSISHQLAAGSLDQKHLRGACACTC 119

OY 113 --GNNLVNKKPERGFTPLIMASAFGEIETVRPLFMGADPHILAKEREASLSASTGYT 170

DB 120 LSGNNLVNKKPERGFTPLIMASAFGEIETVRPLFMGADPHILAKEREASLSASTGYT 179

OY 171 DIVGLLEKRDVVDINIDWNGGTPLLYAVRGHVKCEALLARAGDLTTADSGYTPMDLA 230

DB 180 DIVGLLEKRDVVDINIDWNGGTPLLYAVRGHVKCEALLARAGDLTTADSGYTPMDLA 239

OY 231 VALGYRKYQOVVIEHNLKLFQSNLVPADPE 260

DB 240 VALGYRKYQOVVIEHNLKLFQSNLVPADPE 269

RESULT 4

ID ABP41881 standard; Protein; 229 AA.

AC ABP41881;

DT 22-AUG-2002 (first entry)

DE Human ovarian antigen HWHKD22, SEQ ID NO:3013.

KM Human; ovarian antigen; ovary; breast; cancer; tumour;
 KM ovarian cancer; breast cancer; tumour; reproductive system disorder;
 KM infertility; pregnancy disorder; anovulation; polycystic ovary syndrome;
 KM PCOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection;
 KM inflammatory condition; immune disorder; blood disorder;
 KM cardiovascular disorder; respiratory disorder; neurological disorder;
 KM gastrointestinal disorder; urinary system disorder; drug screening;
 KM gene therapy; chromosome mapping; forensic analysis;
 KM antibody preparation; cytostatic; immunomodulatory; neuroprotective;
 KM antiinflammatory; gynaecological; reproductive.

OS Homo sapiens.

PN WO200200677-A1.

PF 03-JAN-2002.

PR 07-JUN-2001; 2001WO-US18569.

PR 07-JUN-2000; 2000US-209467P.

(HUMA-) HUMAN GENOME SCI INC.

PI Birse CE, Rosen CA;

WPI: 2002-147878/19.

DR N-PSDB; ABQ54958.

PT Isolated nucleic acid molecules encoding novel ovarian polypeptides,
 PT useful in the prevention, treatment and diagnosis of cancer (e.g.
 PT ovarian cancer), immune disorders, cardiovascular disorders and
 PT neurological diseases -

PS Claim 11; SEQ ID NO 3013; 2922pp; English.

CC The invention relates to 2175 novel human ovarian antigens (ABP41054-
 CC ABP43228) and to cDNAs encoding them (ABQ54131-ABQ56305), and also
 CC encompasses polypeptides 90% identical and polynucleotides 95% identical
 CC to the sequences of the invention. The invention additionally relates to
 CC recombinant vectors and host cells comprising human ovarian antigen
 CC polynucleotides, antibodies against human ovarian antigens, and the use
 CC of ovarian antigen polynucleotides and polypeptides in diagnosing,
 CC treating, prognosing or preventing various ovary and/or breast-related
 CC disorders. Such conditions include ovarian cancer and breast cancer, and
 CC metastatic tumours of ovarian or breast origin, reproductive system
 CC disorders (e.g., infertility, disorders of pregnancy, anovulation,
 CC polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine
 CC disorders, infections (e.g., chlamydia, HIV, toxoplasmosis, and toxic
 CC shock syndrome), inflammatory conditions (e.g., mastitis, oophoritis and
 CC vaginitis), immune disorders (e.g., congenital and acquired
 CC immunodeficiencies, autoimmune oophoritis, systemic lupus erythematosus),
 CC blood-related disorders (e.g., anaemia), cardiovascular disorders,
 CC respiratory disorders, neurological disorders, gastrointestinal disorders
 CC and urinary system disorders. Ovarian antigen polypeptides and
 CC polynucleotides may also be used in screening for compounds which

CC modulate ovarian antigen expression or activity. The polynucleotides may
 CC further be used for gene therapy, chromosome mapping, in the
 CC identification of individuals and in forensic analysis, and the
 CC polypeptides may be used as food additives or to prepare antibodies
 CC useful in disease diagnosis, drug targeting and phenotyping. The present
 CC sequence represents a human ovarian antigen of the invention.
 CC Note: The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.

CC Sequence 229 AA;

Query Match 46.6%; Score 625; DB 23; Length 229;
 Best Local Similarity 58.6%; Pred. No. 2.2e-55;
 Matches 130; Conservative 26; Mismatches 58; Indels 8; Gaps 3;

QY 37 TVVLSTPCTPEVNPEDASV-----SSPOAGSLKSTTLTNROGNEVSALPATID 90
 DB 7 TVVFLAECHNIH-TSPSPGIQVRHVYTPSTKHSPIKOSTTLTNKRGNEVSTTPLL 65
 QY 91 SLSTHQLAAQAGELDOLKEHLRKGDNLVKNKPDERTPLIMASAFGEIETVFLLEW 150
 DB 66 SLSTVQLAAQAGELVLAIRIEQ-ENVINHTDEEGFTPLMAAHQIAVEFLQNGADP 124
 QY 151 HLAKERESALSLASTGGYTDIVGLLEERVDINITYMNGGTPLLIYAVRGNHVKCE 210
 DB 125 QLLGKGRESALSLASTGGYTDIVKMLDCCGVADVEYDMNGGTPLLIYAVGHNHVK 184
 QY 211 ARGADLTTEADSGYTPMDLAVALGYRKQOVIENHILKLFOS 252
 DB 185 ESGADPTITDSCGYNSMDLAVALGYSVOVIESHLKLLON 226

RESULT 5
 AAG66309
 ID AAG66309 standard; Protein; 313 AA.

AC AAG66309;

DT 09-OCT-2001 (first entry)

DE Human ankyrin-like protein 34.

KW Human; ankyrin-like protein 34; cytosolic; virucidal; immunomodulatory;
 KW antiinflammatory; haemostatic; gene therapy; malignant tumour;
 KW haemopathy; HIV infection; immunological disease; inflammation.

OS Homo sapiens.

PN WO200155194-A1.

PD 02-AUG-2001.

PF 21-JAN-2001; 2001WO-CN00085.

PR 28-JAN-2000; 2000CN-011595.

PA (BIOD-) BIODOOR GENE TECHNOLOGY LTD SHANGHAI.

PI Mao Y, Xie Y;

DR WPI; 2001-483222/52.

DR N-PSDB; AAH75702.

PT New human ankyrin-like protein 34 for diagnosing and treating malignant
 PT tumor, hemopathy, human immunodeficiency virus infection, immunological
 PT diseases and various inflammations -

PS Claim 1; Page 31-32; 38pp; Chinese.

XX The present sequence is the protein sequence for human ankyrin-like
 CC protein 34. The ankyrin-like protein and its coding sequence are useful
 CC in the diagnosis and treatment of malignant tumour, haemopathy, HIV

CC infection, immunological diseases and various inflammations.

XX Sequence 313 AA;

Query Match 46.3%; Score 621.5; DB 22; Length 313;
 Best Local Similarity 60.6%; Pred. No. 8.1e-55;
 Matches 126; Conservative 26; Mismatches 49; Indels 7; Gaps 2;

QY 51 NPPEDASV-----SSPOAGSLKSTTLTNROGNEVSALPATIDSLIHQAQGEID 104
 DB 104 SPSPGIQVRHVYTPSTKHSPIKOSTTLTNKRGNEVSTTPLLANSUSVHQLAAQAGEL 163
 QY 105 QLKEHLRKGDNLVKNKPDERTPLIMASAFGEIETVFLLEWADPHILAKERESALSLA 164
 DB 164 YLAIRIEQ-ENVINHTDEEGFTPLMAAHQIAVEFLQNGADPOLLGKRESALSLA 222
 QY 165 STGGYTDIVGLLEERVDINITYMNGGTPLLIYAVRGNHVKCEALLARGADLTTEADSGY 224
 DB 223 CSKGYTDIVKMLDCCGVADVEYDMNGGTPLLIYAVGHNHVKCVMLLBSGADPTIETDSGY 282
 QY 225 TPMDLAVALGYRKQOVIENHILKLFOS 252
 DB 283 NSMDLAVALGYSVOVIESHLKLLON 310

RESULT 6
 AAB94322
 ID AAB94322 standard; Protein; 313 AA.

AC AAB94322;

DT 26-JUN-2001 (first entry)

DE Human protein sequence SEQ ID NO:14803.

KW Human; primer; detection; diagnosis; antisense therapy; gene therapy.

OS Homo sapiens.

PN EP1074617-A2.

PD 07-FEB-2001.

PF 28-JUL-2000; 2000EP-0116126.

PR 29-JUL-1999; 99JP-0248036.

PR 27-AUG-1999; 99JP-0300253.

PR 11-JAN-2000; 2000JP-0118776.

PR 02-MAY-2000; 2000JP-0183767.

PR 09-JUN-2000; 2000JP-0241899.

PA (HELI-) HELIX RES INST.

PI Ota T, Isogai T, Nishikawa T, Hayaishi K, Saito K, Yamamoto J;

PI Ishii S, Sugiyama T, Wakamatsu A, Nagai K, Otsuki T;

DR WPI; 2001-318749/34.

PT Primer sets for synthesizing polynucleotides, particularly the 5602

PT full-length cDNAs defined in the specification, and for the detection

PT and/or diagnosis of the abnormality of the proteins encoded by the

PT full-length cDNAs -

PS Claim 8; SEQ ID 14803; 2537pp + CD ROM; English.

XX The present invention describes primer sets for synthesizing 5602
 CC full-length cDNAs defined in the specification. Where a primer set
 CC comprises: (a) an oligo-dT primer and an oligonucleotide complementary
 CC to the complementary strand of a polynucleotide which comprises one of
 CC the 5602 nucleotide sequences defined in the specification, where the
 CC oligonucleotide comprises at least 15 nucleotides; or (b) a combination
 CC of an oligonucleotide comprising a sequence complementary to the
 CC complementary strand of a polynucleotide which comprises a 5'-end

XX

Human; transcription factor; TRFX; cell proliferative disease;

KW developmental disorder; cancer; AIDS; infection; cytostatic; anti-HIV;
 KW neuroprotective; antiinflammatory; gene therapy.
 XX Homo sapiens.
 OS
 XX WO200172777-A2.
 PN
 XX 04-OCT-2001.
 PD
 XX 13-MAR-2001; 2001WO-US08117.
 PF
 XX 13-MAR-2000; 2000US-0188986.
 PR
 XX (INCY-) INCYTE GENOMICS INC.
 PA
 XX Hillman JL, Baughn MR, Yue H, Lai P, Lu DM, Patterson C;
 PI Azimzai Y, Bandman O, Tang YT, Mathur P, Shah P, Au-Young J;
 PI Reddy R;
 XX WPI; 2001-570896/64.
 DR N-PSDB; ABA82985.
 XX
 PT Novel transcription factor polypeptides, used to treat diseases
 associated with altered activity and expression of TRFX, and to screen
 for agents capable of modulating its activity -
 PS
 XX Claim 1; Pages 151-152; 327pp; English.
 CC The present sequence is the protein sequence for a human transcription
 factor. The transcription factor and its coding sequence are useful in
 the diagnosis, treatment and prevention of diseases associated with
 CC altered expression of the transcription factor e.g. cell proliferative,
 CC autoimmune/inflammatory, neurological and developmental disorders. A
 CC number of specific disorders/diseases are given in the specification,
 CC including: arteriosclerosis, cirrhosis, hepatitis, cancers, AIDS,
 CC allergies, anaemia, asthma, autoimmune thyroiditis, bronchitis, atopic
 CC dermatitis, diabetes mellitus, emphysema, Goodpasture's syndrome, gout,
 CC Grave's disease, multiple sclerosis, osteoarthritis, pancreatitis,
 CC psoriasis, rheumatoid arthritis, systemic lupus erythematosus, ulcerative
 CC colitis, uveitis, Alzheimer's disease, Huntington's disease, Parkinson's
 CC disease, stroke, and viral, bacterial, fungal and protozoal infections.
 XX
 SQ Sequence 152 AA;
 Query Match 36.9%; Score 495; DB 22; Length 152;
 Best Local Similarity 68.3%; Pred. No. 2.3e-42;
 Matches 95; Conservative 16; Mismatches 28; Indels 0; Gaps 0;
 QY 114 DNLVKKPDRGFPLIWSAFGEIETVRFLEMGADPHILAKERESALSLASTGTYDIV 173
 Db 11 EVVINTDEEGFPLMAAHGQIAVEFLLQNGADPQLGKRESALSLASTGTYDIV 70
 QY 174 GLLERDVADINIVDNGGTPLLYAVRGNHYKCVBALLAGADLTTEADSGTMDLAVNL 233
 Db 71 KMLLDGCVVNERDMNGGTPLLYAVHGNHYKCVKMLLESADPTLETDSGYNSMDLAVNL 130
 QY 234 GYRKYQGVVIEHNLKLFQS 252
 Db 131 GYRSVQGVIESHLKLQN 149
 RESULT 9
 AAG01584
 ID AAG01584 standard; Protein; 84 AA.
 AC AAG01584;
 XX
 XX 06-OCT-2000 (first entry)
 DT
 XX Human secreted protein, SEQ ID NO: 5665.
 DE
 XX Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation;
 KW gene therapy; chromosome mapping.

XX Homo sapiens.
 OS
 XX EPI033401-A2.
 PN
 XX 06-SEP-2000.
 PD
 XX 21-FEB-2000; 2000EP-0200610.
 PF
 XX 26-FEB-1999; 99US-0122487.
 PR
 XX (GEST) GENSET.
 PA
 XX Dumas Milne Edwards J, Duclert A, Giordano J;
 PI WPI; 2000-500381/45.
 DR N-PSDB; AAC01590.
 XX
 PT New nucleic acid that is a 5' expressed sequence tag (5' EST) for
 obtaining cDNAs and genomic DNAs that correspond to 5' ESTs and for
 PT diagnostic, forensic, gene therapy and chromosome mapping procedures -
 PS
 XX Claim 13; SEQ ID 5665; 71pp + CD-ROM; English.
 CC The present sequence is a polypeptide encoded by one of a large number
 of 5' ESTs derived from mRNAs encoding secreted proteins. The 5' ESTs
 CC were prepared from total human RNAs or polyA+ RNAs derived from 30
 CC different tissues. EST sequences usually correspond mainly to the 3'
 CC untranslated region (UTR) of the mRNA because they are often obtained
 CC from oligo-dT primed cDNA libraries. Such ESTs are not well suited for
 CC isolating cDNA sequences derived from the 5' ends of mRNAs and even in
 CC those cases where longer cDNA sequences have been obtained, the full 5'
 CC UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5'
 CC ends and can therefore be used to obtain full length cDNAs and genomic
 CC DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and
 CC chromosome mapping procedures. They are used to obtain upstream
 CC regulatory sequences and to design expression and secretion vectors.
 XX
 SQ Sequence 84 AA;
 Query Match 31.4%; Score 421.5; DB 21; Length 84;
 Best Local Similarity 98.8%; Pred. No. 3e-35; Indels 1; Gaps 1;
 Matches 84; Conservative 0; Mismatches 0;
 QY 1 MELTQPAEDLIQTQTPASLGDPEDEENADGSDTVVLSLPCTPEPPVNEPDASVS 60
 Db 1 MELTQPAEDLIQTQTPASLGDPEDEENADGSDTVVLSLPCTPEPPVNEPDASVS 60
 QY 61 PQAGSLKHSITLTNRGNEVSAL 85
 Db 61 PQ-GSSLKHSITLTNRGNEVSAL 84
 RESULT 10
 AAU20665
 ID AAU20665 standard; Protein; 119 AA.
 AC AAU20665;
 XX
 XX 04-DEC-2001 (first entry)
 DT
 XX Human secreted protein, Seq ID No 657.
 DE
 XX Immunomodulatory; human immunodeficiency virus; HIV; anaemia; angina;
 KW rheumatoid arthritis; antiarteriosclerotic; cardiant; vascular;
 KW cerebroprotective; thrombolytic; antimicrobial; ophthalmological;
 KW cytotactic; Alzheimer's disease; Parkinson's disease; human; cancer;
 KW multiple sclerosis; cancer; hyperproliferative disorder; infection;
 KW Gaucher's disease; neurological disease; cerebrovascular disorder;
 KW thrombosis; wound healing.
 XX
 OS Homo sapiens.


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RESULT 14
ABB43550
ID ABB43550 standard; Peptide; 49 AA.
XX
AC ABB43550;
XX
DT 04-FEB-2002 (first entry)
XX
DE Peptide #11056 encoded by human foetal liver single exon probe.
XX
KW Human; foetal liver; gene expression; single exon nucleic acid probe.
XX
OS Homo sapiens.
XX
PN WO200157277-A2.
XX
PD 09-AUG-2001.
XX
PF 30-JAN-2001; 2001WO-US00669.
XX
PR 04-FEB-2000; 2000US-0180312.
XX
PR 26-MAY-2000; 2000US-0207456.
XX
PR 30-JUN-2000; 2000US-0608408.
XX
PR 03-AUG-2000; 2000US-0632366.
XX
PR 21-SEP-2000; 2000US-0234687.
XX
PR 27-SEP-2000; 2000US-0236359.
XX
PR 04-OCT-2000; 2000GB-0024263.
XX
PA (MOLE-) MOLECULAR DYNAMICS INC.
XX
PI Penn SG, Hanzel DK, Chen W, Rank DR;
XX
DR WPI; 2001-483447/52.
XX
PT Human genome-derived single exon nucleic acid probes useful for
XX
PT analyzing gene expression in human fetal liver -
XX
PS Claim 27; SEQ ID NO 36185; 639pp + sequence listing; English.
XX
CC The invention relates to a single exon nucleic acid probe for
CC measuring human gene expression in a sample derived from human foetal
CC liver. The single exon nucleic acid probes may be used for predicting,
CC measuring and displaying gene expression in samples derived from human
CC fetal liver. The present sequence is a peptide encoded by a single exon
CC nucleic acid probe of the invention.
CC Note: The sequence data for this patent did not form part of the
CC printed specification, but was obtained in electronic format directly
CC from WIPO at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 49 AA;
XX
Query March 19.2%; Score 257; DB 22; Length 49;
Best Local Similarity 100.0%; Pred. No. 8.5e-19;
Matches 49; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
XX
QY 1 MELTQPAEDLIQTOTPASELGDPEDPGEAAGSDTVVLSLFPCTPP 49
Db 1 MELTQPAEDLIQTOTPASELGDPEDPGEAAGSDTVVLSLFPCTPP 49
XX
RESULT 15
AAM21225
ID AAM21225 standard; Protein; 49 AA.
XX
AC AAM21225;
XX
DT 12-OCT-2001 (first entry)
XX
DE Peptide #7659 encoded by probe for measuring cervical gene expression.
XX
KW Probe; human; microarray; gene expression; cervical epithelial cell;
XX
KW cervical cancer.
XX
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OS Homo sapiens.
XX
PN WO200157278-A2.
XX
PD 09-AUG-2001.
XX
PF 30-JAN-2001; 2001WO-US00670.
XX
PR 04-FEB-2000; 2000US-0180312.
XX
PR 26-MAY-2000; 2000US-0207456.
XX
PR 30-JUN-2000; 2000US-0608408.
XX
PR 03-AUG-2000; 2000US-0632366.
XX
PR 21-SEP-2000; 2000US-0234687.
XX
PR 27-SEP-2000; 2000US-0236359.
XX
PR 04-OCT-2000; 2000GB-0024263.
XX
PA (MOLE-) MOLECULAR DYNAMICS INC.
XX
PI Penn SG, Hanzel DK, Chen W, Rank DR;
XX
DR WPI; 2001-486901/53.
XX
PT Human genome-derived single exon nucleic acid probes useful for
XX
PT analyzing gene expression in human cervical epithelial cells -
XX
PS Claim 27; SEQ ID No 26051; 487pp; English.
XX
CC The present invention relates to human single exon nucleic acid probes
CC (SENPs: see A110068-A128459). The present sequence is a peptide encoded
CC by one such probe. The SENPs are derived from human HeLa cells. The SENPs
CC can be used to produce a single exon microarray, which can be used for
CC measuring human gene expression in a sample derived from human cervical
CC epithelial cells. By measuring gene expression, the probes are therefore
CC useful in grading and/or staging of diseases of the cervix, notably
CC cervical cancer.
CC Note: The sequence data for this patent did not form part of the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ Sequence 49 AA;
XX
Query March 19.2%; Score 257; DB 22; Length 49;
Best Local Similarity 100.0%; Pred. No. 8.5e-19;
Matches 49; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
XX
QY 1 MELTQPAEDLIQTOTPASELGDPEDPGEAAGSDTVVLSLFPCTPP 49
Db 1 MELTQPAEDLIQTOTPASELGDPEDPGEAAGSDTVVLSLFPCTPP 49
XX
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Search completed: March 17, 2003, 16:39:38
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